

Sequence Listing

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<120> METHOD FOR MAKING MONOCLONAL ANTIBODIES AND
CROSS-REACTIVE ANTIBODIES OBTAINABLE BY THE METHOD

<130> P1468R1 (REVISED)

<140> US 09/329,633

<141> 1999-06-10

<150> US 60/089,253

<151> 1998-06-12

<160> 2

<210> 1

<211> 1799

<212> DNA

<213> human

<400> 1

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ccacgggcct gagagactat aagagcgttc cctaccgcca tggaacaacg 150
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cacgatgctg ataaagtggg tcaacaaaac cgggcgagat gcctctgtcc 1250
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agtaggaaag tgccacaatt gtcacatgac cgggtactgga agaaactctc 1500
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gtctggatca ttccgtttgt gcgtactttg agatttggtt tgggatgtca 1650
ttgttttcac agcacttttt taccctaagc taaatgcttt atttatttat 1700
ttgggctaca ttgtaagatc catctacaaa aaaaaaaaaa aaaaaaaag 1750
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<210> 2
<211> 411
<212> PRT
<213> human

<220>

<221> xaa

<222> 410

<223> xaa = leu or met

<400> 2

Met	Glu	Gln	Arg	Gly	Gln	Asn	Ala	Pro	Ala	Ala	Ser	Gly	Ala	Arg
1				5				10					15	

Lys	Arg	His	Gly	Pro	Gly	Pro	Arg	Glu	Ala	Arg	Gly	Ala	Arg	Pro
			20					25					30	

Gly	Leu	Arg	Val	Pro	Lys	Thr	Leu	Val	Leu	Val	Val	Ala	Ala	Val
			35					40					45	

Leu	Leu	Leu	Val	Ser	Ala	Glu	Ser	Ala	Leu	Ile	Thr	Gln	Gln	Asp
			50					55					60	

Leu	Ala	Pro	Gln	Gln	Arg	Ala	Ala	Pro	Gln	Gln	Lys	Arg	Ser	Ser
			65					70					75	

Pro	Ser	Glu	Gly	Leu	Cys	Pro	Pro	Gly	His	His	Ile	Ser	Glu	Asp
			80					85					90	

Gly	Arg	Asp	Cys	Ile	Ser	Cys	Lys	Tyr	Gly	Gln	Asp	Tyr	Ser	Thr
			95					100					105	

His	Trp	Asn	Asp	Leu	Leu	Phe	Cys	Leu	Arg	Cys	Thr	Arg	Cys	Asp
			110					115					120	

Ser	Gly	Glu	Val	Glu	Leu	Ser	Pro	Cys	Thr	Thr	Thr	Arg	Asn	Thr
			125					130					135	

Val	Cys	Gln	Cys	Glu	Glu	Gly	Thr	Phe	Arg	Glu	Glu	Asp	Ser	Pro
			140					145					150	

Glu	Met	Cys	Arg	Lys	Cys	Arg	Thr	Gly	Cys	Pro	Arg	Gly	Met	Val
			155					160					165	

Lys	Val	Gly	Asp	Cys	Thr	Pro	Trp	Ser	Asp	Ile	Glu	Cys	Val	His
			170					175					180	

Lys	Glu	Ser	Gly	Ile	Ile	Ile	Gly	Val	Thr	Val	Ala	Ala	Val	Val
			185					190					195	

Leu	Ile	Val	Ala	Val	Phe	Val	Cys	Lys	Ser	Leu	Leu	Trp	Lys	Lys
			200					205					210	

Val	Leu	Pro	Tyr	Leu	Lys	Gly	Ile	Cys	Ser	Gly	Gly	Gly	Gly	Asp
			215					220					225	

Pro Glu Arg Val Asp Arg Ser Ser Gln Arg Pro Gly Ala Glu Asp		
230	235	240
Asn Val Leu Asn Glu Ile Val Ser Ile Leu Gln Pro Thr Gln Val		
245	250	255
Pro Glu Gln Glu Met Glu Val Gln Glu Pro Ala Glu Pro Thr Gly		
260	265	270
Val Asn Met Leu Ser Pro Gly Glu Ser Glu His Leu Leu Glu Pro		
275	280	285
Ala Glu Ala Glu Arg Ser Gln Arg Arg Arg Leu Leu Val Pro Ala		
290	295	300
Asn Glu Gly Asp Pro Thr Glu Thr Leu Arg Gln Cys Phe Asp Asp		
305	310	315
Phe Ala Asp Leu Val Pro Phe Asp Ser Trp Glu Pro Leu Met Arg		
320	325	330
Lys Leu Gly Leu Met Asp Asn Glu Ile Lys Val Ala Lys Ala Glu		
335	340	345
Ala Ala Gly His Arg Asp Thr Leu Tyr Thr Met Leu Ile Lys Trp		
350	355	360
Val Asn Lys Thr Gly Arg Asp Ala Ser Val His Thr Leu Leu Asp		
365	370	375
Ala Leu Glu Thr Leu Gly Glu Arg Leu Ala Lys Gln Lys Ile Glu		
380	385	390
Asp His Leu Leu Ser Ser Gly Lys Phe Met Tyr Leu Glu Gly Asn		
395	400	405
Ala Asp Ser Ala Xaa Ser		
410		